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PATENT APPLICATION: US/09/609,027B

DATE: 03/01/2002

TIME: 15:30:49

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\03012002\I609027B.raw

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3 <110> APPLICANT: HENDRICKSON, WAYNE
 4 JIANG, XULIANG
 5 LANGLEY, KEITH E
 6 SYED, RASHID
 7 ANN HSU, YUEH-RONG
 9 <120> TITLE OF INVENTION: CONJUGATED LIGANDS FOR THE STIMULATION OF BLOOD CELL
 PROLIFERATION BY
 10 EFFECTING DIMERIZATION OF THE RECEPTOR FOR STEM CELL FACTOR
 12 <130> FILE REFERENCE: 0575/50950
 14 <140> CURRENT APPLICATION NUMBER: US 09/609,027B
 C--> 15 <141> CURRENT FILING DATE: 2000-06-29
 17 <160> NUMBER OF SEQ ID NOS: 22
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 141
 23 <212> TYPE: PRT
 24 <213> ORGANISM: HOMO SAPIENS
 26 <400> SEQUENCE: 1
 28 Glu Gly Ile Cys Arg Asn Arg Val Thr Asn Asn Val Lys Asp Val Thr
 29 1 5 10 15
 32 Lys Leu Val Ala Asn Leu Pro Lys Asp Tyr Met Ile Thr Leu Lys Tyr
 33 20 25 30
 36 Val Pro Gly Met Asp Val Leu Pro Ser His Gln Trp Ile Ser Glu Met
 37 35 40 45
 40 Val Val Gln Leu Ser Asp Ser Leu Thr Asp Leu Leu Asp Lys Phe Ser
 41 50 55 60
 44 Asn Ile Ser Glu Gly Leu Ser Asn Tyr Ser Ile Ile Asp Lys Leu Val
 45 65 70 75 80
 48 Asn Ile Val Asp Asp Leu Val Glu Cys Val Lys Glu Asn Ser Ser Lys
 49 85 90 95
 52 Asp Leu Lys Lys Ser Phe Lys Ser Pro Glu Pro Arg Leu Phe Thr Pro
 53 100 105 110
 56 Glu Glu Phe Phe Arg Ile Phe Asn Arg Ser Ile Asp Ala Phe Lys Asp
 57 115 120 125
 60 Phe Val Val Ala Ser Glu Thr Ser Asp Cys Val Val Ser
 61 130 135 140
 64 <210> SEQ ID NO: 2
 65 <211> LENGTH: 150
 66 <212> TYPE: PRT
 67 <213> ORGANISM: HOMO SAPIENS
 69 <400> SEQUENCE: 2
 71 Glu Glu Val Ser Glu Tyr Cys Ser His Met Ile Gly Ser Gly His Leu
 72 1 5 10 15
 75 Gln Ser Leu Gln Arg Leu Ile Asp Ser Gln Met Glu Thr Ser Cys Gln

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76          20          25          30
79 Ile Thr Phe Glu Phe Val Asp Gln Glu Gln Leu Lys Asp Pro Val Cys
80          35          40          45
83 Tyr Leu Lys Lys Ala Phe Leu Leu Val Gln Asp Ile Met Glu Asp Thr
84          50          55          60
87 Met Arg Phe Arg Asp Asn Thr Pro Asn Ala Ile Ala Ile Val Gln Leu
88 65          70          75          80
91 Gln Glu Leu Ser Leu Arg Leu Lys Ser Cys Phe Thr Lys Asp Tyr Glu
92          85          90          95
95 Glu His Asp Lys Ala Cys Val Arg Thr Phe Tyr Glu Thr Pro Leu Gln
96          100          105          110
99 Leu Leu Glu Lys Val Lys Asn Val Phe Asn Glu Thr Lys Asn Leu Leu
100          115          120          125
103 Asp Lys Asp Trp Asn Ile Phe Ser Lys Asn Cys Asn Asn Ser Phe Ala
104          130          135          140
107 Glu Cys Ser Ser Gln Gly
108 145          150
111 <210> SEQ ID NO: 3
112 <211> LENGTH: 129
113 <212> TYPE: PRT
114 <213> ORGANISM: HOMO SAPIENS
116 <400> SEQUENCE: 3
118 His Lys Cys Asp Ile Thr Leu Gln Glu Ile Ile Lys Thr Leu Asn Ser
119 1          5          10          15
122 Leu Thr Glu Gln Lys Thr Leu Cys Thr Glu Leu Thr Val Thr Asp Ile
123          20          25          30
126 Phe Ala Ala Ser Lys Asn Thr Thr Glu Lys Glu Thr Phe Cys Arg Ala
127          35          40          45
130 Ala Thr Val Leu Arg Gln Phe Tyr Ser His His Glu Lys Asp Thr Arg
131          50          55          60
134 Cys Leu Gly Ala Thr Ala Gln Gln Phe His Arg His Lys Gln Leu Ile
135 65          70          75          80
138 Arg Phe Leu Lys Arg Leu Asp Arg Asn Leu Trp Gly Leu Ala Gly Leu
139          85          90          95
142 Asn Ser Cys Pro Val Lys Glu Ala Asn Gln Ser Thr Leu Glu Asn Phe
143          100          105          110
146 Leu Glu Arg Leu Lys Thr Ile Met Arg Glu Lys Tyr Ser Lys Cys Ser
147          115          120          125
150 Ser
154 <210> SEQ ID NO: 4
155 <211> LENGTH: 127
156 <212> TYPE: PRT
157 <213> ORGANISM: HOMO SAPIENS
159 <400> SEQUENCE: 4
161 Ala Pro Ala Arg Ser Pro Ser Pro Ser Thr Gln Pro Trp Glu His Val
162 1          5          10          15
165 Asn Ala Ile Gln Glu Ala Arg Arg Leu Leu Asn Leu Ser Arg Asp Thr
166          20          25          30
169 Ala Ala Glu Met Asn Glu Thr Val Glu Val Ile Ser Glu Met Phe Asp

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170          35          40          45
173 Leu Gln Glu Pro Thr Cys Leu Gln Thr Arg Leu Glu Leu Tyr Lys Gln
174          50          55          60
177 Gly Leu Arg Gly Ser Leu Thr Lys Ile Lys Gly Pro Leu Thr Met Met
178 65          70          75          80
181 Ala Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro Glu Thr Ser Cys
182          85          90          95
185 Ala Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu Asn Leu Lys Asp
186          100          105          110
189 Phe Leu Leu Val Ile Pro Phe Asp Cys Trp Glu Pro Val Gln Glu
190          115          120          125
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 132
195 <212> TYPE: PRT
196 <213> ORGANISM: HOMO SAPIENS
198 <400> SEQUENCE: 5
200 Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln Leu Gln Leu Glu His
201 1          5          10          15
204 Leu Leu Leu Asp Leu Gln Met Ile Leu Asn Gly Ile Asn Asn Tyr Lys
205          20          25          30
208 Asn Pro Lys Leu Thr Arg Met Leu Thr Phe Lys Tyr Met Pro Lys Lys
209          35          40          45
212 Ala Thr Glu Leu Lys His Leu Gln Cys Leu Glu Glu Glu Leu Lys Pro
213          50          55          60
216 Leu Glu Glu Val Leu Asn Leu Ala Gln Ser Lys Asn Phe His Leu Arg
217 65          70          75          80
220 Pro Arg Asp Leu Ile Ser Asn Ile Asn Val Ile Val Leu Glu Leu Lys
221          85          90          95
224 Gly Ser Glu Thr Thr Phe Met Cys Glu Tyr Ala Asp Glu Thr Ala Thr
225          100          105          110
228 Ile Val Glu Phe Leu Asn Arg Trp Ile Thr Phe Cys Gln Ser Ile Ile
229          115          120          125
232 Ser Thr Leu Thr
233          130
236 <210> SEQ ID NO: 6
237 <211> LENGTH: 115
238 <212> TYPE: PRT
239 <213> ORGANISM: HOMO SAPIENS
241 <400> SEQUENCE: 6
243 Ile Pro Thr Glu Ile Pro Thr Ser Ala Leu Val Lys Glu Thr Leu Ala
244 1          5          10          15
247 Leu Leu Ser Thr His Arg Thr Leu Leu Ile Ala Asn Glu Thr Leu Arg
248          20          25          30
251 Ile Pro Val Pro Val His Lys Asn His Gln Leu Cys Thr Glu Glu Ile
252          35          40          45
255 Phe Gln Gly Ile Gly Thr Leu Glu Ser Gln Thr Val Gln Gly Gly Thr
256          50          55          60
259 Val Glu Arg Leu Phe Lys Asn Leu Ser Leu Ile Lys Lys Tyr Ile Asp
260 65          70          75          80

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263 Gly Gln Lys Lys Lys Cys Gly Glu Glu Arg Arg Arg Val Asn Gln Phe
264                               85                               90           95
267 Leu Asp Tyr Leu Gln Glu Phe Leu Gly Val Met Asn Thr Glu Trp Ile
268                               100                               105           110
271 Ile Glu Ser
272                               115
275 <210> SEQ ID NO: 7
276 <211> LENGTH: 164
277 <212> TYPE: PRT
278 <213> ORGANISM: RATTUS SP.
280 <400> SEQUENCE: 7
282 Glu Gly Ile Cys Arg Asn Arg Val Thr Asn Asn Val Lys Asp Val Thr
283 1                               5                               10           15
286 Lys Leu Val Ala Asn Leu Pro Lys Asp Tyr Met Ile Thr Leu Lys Tyr
287                               20                               25           30
290 Val Pro Gly Met Asp Val Leu Pro Ser His Cys Trp Ile Ser Glu Met
291                               35                               40           45
294 Val Val Gln Leu Ser Asp Ser Leu Thr Asp Leu Leu Asp Lys Phe Ser
295                               50                               55           60
298 Asn Ile Ser Glu Gly Leu Ser Asn Tyr Ser Ile Ile Asp Lys Leu Val
299 65                               70                               75           80
302 Asn Ile Val Asp Asp Leu Val Glu Cys Val Lys Glu Asn Ser Ser Lys
303                               85                               90           95
306 Asp Leu Lys Lys Ser Phe Lys Ser Pro Glu Pro Arg Leu Phe Thr Pro
307                               100                               105           110
310 Glu Glu Phe Phe Arg Ile Phe Asn Arg Ser Ile Asp Ala Phe Lys Asp
311                               115                               120           125
314 Phe Val Val Ala Ser Glu Thr Ser Asp Cys Val Val Ser Ser Thr Leu
315                               130                               135           140
318 Ser Pro Glu Lys Asp Ser Arg Val Ser Val Thr Lys Pro Phe Met Leu
319 145                               150                               155           160
322 Pro Pro Val Ala
326 <210> SEQ ID NO: 8
327 <211> LENGTH: 164
328 <212> TYPE: PRT
329 <213> ORGANISM: MURINE SP.
331 <400> SEQUENCE: 8
333 Lys Glu Ile Cys Gly Asn Pro Val Thr Asp Asn Val Lys Asp Ile Thr
334 1                               5                               10           15
337 Lys Leu Val Ala Asn Leu Pro Asn Asp Tyr Met Ile Thr Leu Asn Tyr
338                               20                               25           30
341 Val Ala Gly Met Asp Val Leu Pro Ser His Cys Trp Leu Arg Asp Met
342                               35                               40           45
345 Val Ile Gln Leu Ser Leu Ser Leu Thr Thr Leu Leu Asp Lys Phe Ser
346                               50                               55           60
349 Asn Ile Ser Glu Gly Leu Ser Asn Tyr Ser Ile Ile Asp Lys Leu Gly
350 65                               70                               75           80
353 Lys Ile Val Asp Asp Leu Val Leu Cys Met Glu Glu Asn Ala Pro Lys
354                               85                               90           95

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357 Asn Ile Lys Glu Ser Pro Lys Arg Pro Glu Thr Arg Ser Phe Thr Pro 110
358 100
361 Glu Glu Phe Phe Ser Ile Phe Asn Arg Ser Ile Asp Ala Phe Lys Asp 125
362 115
365 Phe Met Val Ala Ser Asp Thr Ser Asp Cys Val Leu Ser Ser Thr Leu 140
366 130
369 Ser Pro Glu Lys Asp Ser Arg Val Ser Val Thr Lys Pro Phe Met Leu 160
370 145
373 Pro Pro Val Ala
377 <210> SEQ ID NO: 9
378 <211> LENGTH: 164
379 <212> TYPE: PRT
380 <213> ORGANISM: RATTUS SP.
382 <400> SEQUENCE: 9
384 Gln Glu Ile Cys Arg Asn Pro Val Thr Asp Asn Val Lys Asp Ile Thr 15
385 1 5
388 Lys Leu Val Ala Asn Leu Pro Asn Asp Tyr Met Ile Thr Leu Asn Tyr 30
389 20
392 Val Ala Gly Met Asp Val Leu Pro Ser His Cys Trp Leu Arg Asp Met 45
393 35
396 Val Thr His Leu Ser Val Ser Leu Thr Thr Leu Leu Asp Lys Phe Ser 60
397 50
400 Asn Ile Ser Glu Gly Leu Ser Asn Tyr Ser Ile Ile Asp Lys Leu Gly 80
401 65 70
404 Lys Ile Val Asp Asp Leu Val Ala Cys Met Glu Glu Asn Ala Pro Lys 95
405 85
408 Asn Val Lys Glu Ser Leu Lys Lys Pro Glu Thr Arg Asn Phe Thr Pro 110
409 100
412 Glu Glu Phe Phe Ser Ile Phe Asn Arg Ser Ile Asp Ala Phe Lys Asp 125
413 115
416 Phe Met Val Ala Ser Asp Thr Ser Asp Cys Val Leu Ser Ser Thr Leu 140
417 130
420 Ser Pro Glu Lys Asp Ser Arg Val Ser Val Thr Lys Pro Phe Met Leu 160
421 145
424 Pro Pro Val Ala
428 <210> SEQ ID NO: 10
429 <211> LENGTH: 165
430 <212> TYPE: PRT
431 <213> ORGANISM: CANIS SP.
433 <400> SEQUENCE: 10
435 Lys Gly Ile Cys Gly Lys Arg Val Thr Asp Asn Val Lys Asp Val Thr 15
436 1 5
439 Lys Leu Val Ala Asn Leu Pro Lys Asp Tyr Lys Ile Ala Leu Lys Tyr 30
440 20
443 Val Pro Gly Met Asp Val Leu Pro Ser His Cys Trp Ile Ser Val Met 45
444 35
447 Val Glu Gln Leu Ser Val Ser Leu Thr Asp Leu Leu Asp Lys Phe Ser 60
448 50
451 Asn Ile Ser Glu Gly Leu Ser Asn Tyr Ser Ile Ile Asp Lys Leu Val 160

VERIFICATION SUMMARY

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